## IN THE CLAIMS

Please amend the claims as follows:

Claim 1 (Currently Amended): A method of determining the germination vigour, and/or the storage capability, or the germination vigour and the storage capacity of a seed batch, characterized in that it comprises comprising quantifying, on a sample of seeds taken from said batch, the one or more proteins recognized by at least one anti-L-isoaspartyl methyltransferase antibodies antibody directed against a region of said protein one or more proteins, wherein the region is selected from the group consisting of

a region defined by the sequence: RYVPLTSRX<sub>1</sub>X<sub>2</sub>QLX<sub>3</sub> (SEQ ID NO: 1),

wherein  $X_1$  is E, V or S,  $X_2$  is A or E, and  $X_3$  is R, G or Q, a region defined by the sequence:  $QX_4LX_5VX_6DKX_7X_8DGSX_9X_{10}X_{11}$  (SEQ ID NO: 2).

wherein  $X_4$  is D or E,  $X_5$  is Q or K,  $X_6$  is V or I,  $X_7$  is N or S,  $X_8$  is S, E or A,  $X_9$  is either a dipeptide selected from the group consisting of IS, VS, VT and TS, or a peptide bond,  $X_{10}$  is I or V, and  $X_{11}$  is K, Q or R, a region defined by the sequence: QDLQVVDKNSDGSVSIK (SEQ ID NO: 3), and a region defined by the sequence: RYVPLTSREAQLR (SEQ ID NO: 5)

defined by the sequence (I): RYVPLTSRX<sub>1</sub>X<sub>2</sub>QLX<sub>3</sub> (SEQ ID NO: 1), in which X<sub>1</sub> represents E, V or S, X<sub>2</sub> represents A or E, and X<sub>3</sub> represents R, G or Q.

Claim 2 (Currently Amended): The method as claimed in claim 1, characterized in that the quantification of the L isoaspartyl methyltransferase is carried out using an wherein the region is the region defined by the sequence: RYVPLTSRX<sub>1</sub>X<sub>2</sub>QLX<sub>3</sub> (SEQ ID NO: 1), wherein X<sub>1</sub> is E, V or S, X<sub>2</sub> is A or E, and X<sub>3</sub> is R, G or Q anti L isoaspartyl methyltransferase antibody chosen from:

Claim 4 (Currently Amended): The An anti-L-isoaspartyl methyltransferase antibody as defined in either of claims 2 and 3 selected from the group consisting of

an anti-L-isoaspartyl methyltransferase antibody directed against a protein region defined by the sequence:  $RYVPLTSRX_1X_2QLX_3$  (SEQ ID NO: 1), wherein  $X_1$  is E, V or S,  $X_2$  is A or E, and  $X_3$  is R, G or Q,

an anti-L-isoaspartyl methyltransferase antibody directed against a protein region defined by the sequence: QX<sub>4</sub>LX<sub>5</sub>VX<sub>6</sub>DKX<sub>7</sub>X<sub>8</sub>DGSX<sub>9</sub>X<sub>10</sub>X<sub>11</sub> (SEQ ID NO: 2), wherein X<sub>4</sub> is D or E, X<sub>5</sub> is Q or K, X<sub>6</sub> is V or I, X<sub>7</sub> is N or S, X<sub>8</sub> is S, E or A, X<sub>9</sub> is either a dipeptide selected from the group consisting of IS, VS, VT and TS, or a peptide bond, X<sub>10</sub> is I or V, and X<sub>11</sub> is K, Q or R, an antibody directed against a protein region defined by the sequence:

QDLQVVDKNSDGSVSIK (SEQ ID NO: 3), and
an antibody directed against a protein region defined by the sequence:

RYVPLTSREAQLR (SEQ ID NO: 5).

Claim 5 (Currently Amended): A method of quantifying the L-isoaspartyl methyltransferase in plant material, characterized in that it comprises comprising bringing said <u>plant</u> material into contact with <u>an the</u> anti-L-isoaspartyl methyltransferase antibody as claimed in claim 4.

Claim 6 (Canceled).

Claim 7 (New): The method as claimed in claim 1, wherein the region is the region defined by the sequence: QDLQVVDKNSDGSVSIK (SEQ ID NO: 3).

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Claim 8 (New): The method as claimed in claim 1, wherein the region is the region defined by the sequence: RYVPLTSREAQLR (SEQ ID NO: 5).